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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
09/701,586	11/30/2000	Michael Kock	49100	5846	
7:	590 09/24/2002				
Keil & Weink		EXAMINER			
1101 Connecticut Avenue NW Washington, DC 20036			CHEN, LIPING		
			ART UNIT	PAPER NUMBER	
			1632	1	
			DATE MAILED: 09/24/2002	Ŋ	

Please find below and/or attached an Office communication concerning this application or proceeding.

		Application	No.	Applicant(s)
		09/701,586		KOCK ET AL.
	Office Action Summary	Examiner		Art Unit
		Liping Che	n	1632
Period fo	- The MAILING DATE of this communication app r Reply	pears on the o	cover sheet with the c	orrespondence address
THE N - Exten after S - If the - If NO - Failur - Any re	ORTENED STATUTORY PERIOD FOR REPLY MAILING DATE OF THIS COMMUNICATION. sions of time may be available under the provisions of 37 CFR 1.13 SIX (6) MONTHS from the mailing date of this communication. period for reply specified above is less than thirty (30) days, a reply period for reply is specified above, the maximum statutory period we to reply within the set or extended period for reply will, by statute, exply received by the Office later than three months after the mailing dipatent term adjustment. See 37 CFR 1.704(b).	36(a). In no eventy within the statutowill apply and will apply and will applicate the	t, however, may a reply be time ory minimum of thirty (30) days expire SIX (6) MONTHS from ation to become ABANDONEI	ely filed s will be considered timely. the mailing date of this communication. O (35 U.S.C. § 133).
1)⊠	Responsive to communication(s) filed on 30 N	November 20	000 .	
2a)☐		is action is n		•
3)□	Since this application is in condition for allowardosed in accordance with the practice under			
Disposition	on of Claims			
4)⊠	Claim(s) 1-32 is/are pending in the application	٦.		
4	4a) Of the above claim(s) is/are withdraw	wn from cons	sideration.	
5)	Claim(s) is/are allowed.			
6)	Claim(s) is/are rejected.			
7)	Claim(s) is/are objected to.			
	Claim(s) <u>1-32</u> are subject to restriction and/or e	election requ	irement.	
_ ``	on Papers			
l '	The specification is objected to by the Examine		ti titi butba Fua	i
10)∐ I	The drawing(s) filed on is/are: a) accep			
111	Applicant may not request that any objection to the The proposed drawing correction filed on		·	
'')''	If approved, corrected drawings are required in rep			e
12) 🗆 1	The oath or declaration is objected to by the Ex		oo donon.	
, —	nder 35 U.S.C. §§ 119 and 120			
-	Acknowledgment is made of a claim for foreign	n priority und	er 35 U.S.C. § 119(a)-(d) or (f).
· —	☐ All b)☐ Some * c)☐ None of:		3	, (-, - (,
	1. ☐ Certified copies of the priority document	ts have been	received.	
	2. Certified copies of the priority document			on No
	Copies of the certified copies of the prio application from the International Buse the attached detailed Office action for a list	ority documer ureau (PCT R	nts have been receive Rule 17.2(a)).	ed in this National Stage
	cknowledgment is made of a claim for domesti		·	
a) ☐ The translation of the foreign language pro Acknowledgment is made of a claim for domest	ovisional app	lication has been rec	eived.
Attachmen	_	as priority un	40, 00 0.0.0. 33 120	GIIGIGI TETI
1) Notic	e of References Cited (PTO-892) e of Draftsperson's Patent Drawing Review (PTO-948) nation Disclosure Statement(s) (PTO-1449) Paper No(s)		· ==	/ (PTO-413) Paper No(s) Patent Application (PTO-152)

Art Unit: 1632

Sequence Compliance

This application fails to comply with requirement of 37 CFR 1.821-1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures.

This application refers to general sequence motifs:

- 1) PX_n(S/T)GX₃GKGIYFA, (S/T)XGLR(I/V)XPX_n(S/T)GX₃GKGIYFA
- 2) $LLWHG(S/T)X_7IL(S/T)XGLR(I/V) XPX_n(S/T)GX_3GKGIYFAX_3SKSAXY$
- 3) LX₉NX₂YX₂QLLX(D/E)X_{10/11}WGRVG, AX₃FXKX₄KTXNXWX₅FX₃PXK
- 4) QXL(I/L)X₂IX₉MX₁₀PLGKLX₃QIX₆L, FYTXIPHXFGX₃PP, or KX₃LX₂LXDIEXAX₂L

However, no CRF or paper has been received. Applicant must comply with the requirements of the sequence rules (37 CFR 1.821-1.825).

Election/Restriction

Lack of unity is required under 35 U.S.C. 121 and 372. This application contains the following inventions or goups of inventions which are not so linked as to form a single general inventive concept under PCT Rule 13.1.

- I. Claims 1-4, drawn to a poly(ADP-ribose) polymerase (PARP) homolog containing a functional NAD⁺ binding domain but without zinc finger sequence motif of the general formula.
- II. Claims 5, drawn to a binding partner for PARP homologs, which is an antibody.
- III. Claims 5, drawn to a binding partner for PARP homologs, which is a protein-like compound.

- IV Claims 5, 31 and 32, drawn to a binding partner for PARP homologs, which is a low molecular weight effector, and a method of using the PARP binding partners for the diagnosis of pathological states.
- V. Claims 6-10, drawn to a nucleic acid encoding at least one PARP homolog, an expression cassette, a recombinant vector, and a recombinant microorganism comprising at least one recombinant vector.
- VI. Claim 11, drawn to a transgenic mammal comprising at least one recombinant vector.
- VII. Claim 12, drawn to PARP-deficient eukaryotic cell or a PARP-deficient mammal.
- VIII. Claims 13-22, a method of in vitro detection for PARP inhibitor using PARP homolog.
- IX. Claim 23, drawn to a method for in vitro screening binding partners for a PARP molecule.
- X. Claim 24 and 27, drawn to a method for the qualitative or quantitative determination of nucleic acids encoding a PARP homolog.
- XI. Claim 25 and 26, drawn to a method for the qualitative or quantitative determination of a PARP homolog.
- XII. Claim 28, drawn to a method for determining the efficacy of PAPR effectors.
- XIII. Claim 29, drawn to a gene therapy composition comprising an antisense nucleic acid against a coding nucleic acid which encodes a PARP homolog.
- XIV. Claim 30, drawn to a pharmaceutical composition comprising at least one PARP protein.
- XV. Claims 31 and 32, drawn to a method of using low molecular weight PAPR binding partners for therapy.

This application contains claims directed to more than one invention. These inventions are:

- 1) PX_n(S/T)GX₃GKGIYFA, (S/T)XGLR(I/V)XPX_n(S/T)GX₃GKGIYFA

 LLWHG(S/T)X₇IL(S/T)XGLR(I/V) XPX_n(S/T)GX₃GKGIYFAX₃SKSAXY
- 2) LX₉NX₂YX₂QLLX(D/E)X_{10/11}WGRVG, AX₃FXKX₄KTXNXWX₅FX₃PXK

 QXL(I/L)X₂IX₉MX₁₀PLGKLX₃QIX₆L, FYTXIPHXFGX₃PP, or KX₃LX₂LXDIEXAX₂L

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3) SEQ ID NO:1-10.

These are deemed to lack unity of invention because they are not so linked as to form a single general inventive concept under PCT Rule 13.1. They are different inventions because they are different sequences derived from different genes and have different chemical structures, physical properties and biological functions. Upon election of a group, applicant is required to select one sequence for examination practice.

The inventions listed as Groups I-XV do not relate to a single general inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons: Group I encompasses a PARP homolog and a method of in vitro detection for PARP inhibitor. Group II-XV are directed to different products or methods that require different special technical features as summarized as follows: Group II-VII, XIII and XIV are directed to different products as a PARP binding partner as an antibody, a PARP binding partner as a protein-like compound, a PARP binding partner as a low molecular weight effector, a microorganism comprising a PARP homolog, a transgenic mammal comprising PARP homolog, a PARP-deficient eukaryotic cell or a PARP-deficient mammal, a gene therapy composition comprising an antisense nucleic acid against a nucleic acid encoding PAPR homolog, or a pharmaceutical composition comprising PARP protein, respectively; Groups IX-XII and XV are directed to mutually exclusive methods, each method require specific technique feature other than the technique used in Group I. Moreover, the PARP homolog, which has a functional NAD+ binding domain but no zinc finger sequence motif of the general formula of CX₂CM_mHX₂C is anticipated by Lepiniec et al. (FEBS 364:103-108, 1995). Thus, Groups I-XV lack a common special technical feature. Further, 37 CFR 1.475 does not provide for multiple independent products, methods of manufacture and methods of use (37 CFR 1.475(d). Therefore, The inventions listed as Groups I-XV do not relate to a single general inventive concept under PCT Rule 13.1.

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Applicant is advised that the reply to this requirement to be complete must include an election of

the invention to be examined even though the requirement be traversed (37 CFR 1.143).)

Applicant is reminded that upon the cancellation of claims to a non-elected invention, the

inventorship must be amended in compliance with 37 CFR 1.48(b) if one or more of the currently named

inventors is no longer an inventor of at least one claim remaining in the application. Any amendment of

inventorship must be accompanied by a request under 37 CFR 1.48(b) and by the fee required under 37

CFR 1.17(i).

Any inquiry concerning this communication or earlier communications from the examiner should

be directed to Liping Chen, whose telephone number is (703) 305-4842. The examiner can normally be

reached on Monday through Friday from 8:00 to 5:00 (Eastern Standard Time). Should the examiner be

unavailable, inquiries should be directed to Deborah Reynolds, Supervisory Primary Examiner of Art Unit

1632, at (703) 305-4051. Any administrative or procedural questions should be directed to Pauline

Farrier, Patent Analyst, at (703) 305-2758. Papers related to this application may be submitted to Group

1600 by facsimile transmission. Papers should be faxed to Group 1600 via the PTO Fax Center located in

Crystal Mall 1. The faxing of such papers must conform with the notice published in the Official Gazette,

1096 OG 30 (November 15, 1989). The CM1 Fax Center number is (703) 308-8724.

The Group and/or Art Unit location of your application in the PTO has changed. To aid in

correlating any papers for this application, all further correspondence regarding this application should be

directed to Group Art Unit 1632.

Liping Chen, Ph.D. Patent Examiner

Group 1632

September 5, 2002

PATENT EXAMINER

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Application No.:	
ATENT A LICATIONS CONTAINING	

NOTICE TO COMPLY WITH EQUIREMENTS FOR PATENT AND LICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not	
comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for	the
following reason(s):	

ng r	reason(s):
Ø	 This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
	2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
	3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
	6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
	7. Other:
Apı	plicant Must Provide:
M	An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
<u>D</u>	An initial or <u>substitute</u> paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
Ą	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
For	questions regarding compliance to these requirements, please contact:
For	Rules Interpretation, call (703) 308-4216 CRF Submission Help, call (703) 308-4212 tentIn Software Program Support Technical Assistance703-287-0200
	To Purchase Patentin Software703-306-2600

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY